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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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54:gnEST6 55:gnEST7 56:gnEST3 57:gnEST3 58:gnEST10
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(1-1527) from US08469637A.seq
1527
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ALIGNMENTS

TITLE JOURNAL COMMENT	AUTHORS	ORGANISM REFERENCE	NID KEYWORDS SOURCE	DEFINITION ACCESSION	RESULT 1 LOCUS
The WashU-Merck EST Project Unpublished (1995) Contact: Wilson RK WashU-Merck EST Project Washington University School of Medicine	Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.	Homo sapiens Eukaryotae; mitochondrial eukaryotes; Metazoa; Chordata; Vertebrata; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 463)	g1856711 EST. human.	rl Soares NhHMPu Sl Homo sapiens cDNA clon	AA233719 463 bp mRNA EST 28-FEB-1997

Result

Query

SUMMARIES

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Fax: 314 286 1810
Email: est@watson.wustl.edu
This clone is available
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5,000 clones made from the same 3 libraries. consisted of I.M.A.G.E. clones 260232-265223, 340488-345479, and 484488-489479."
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Key Location/Qualifiers
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Contact: Wilson
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zr47g08.rl Soares
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of I.M.A.G.E.
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/note="Organ: mixed (see below); Vector: pT7T3D-Pac
/note="Organ: mixed (see below); Vector: pT7T3D-Pac
(Pharmacia) with a modified polylinker; Site_1: Not I;
Site_2: Eco RI; Equal amounts of plasmid DNA from three
normalized libraries (melanocyte 2NbHM, pregnant uterus
NbHPU, and fetal heart NbHH19W) were mixed, and ss circles
were made in vitro. Following HAP purification, this DNA
was used as tracer in a subtractive hybridization reaction
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tggatgagtaaactgtttctcaggcacttgaggctttcagtgatatctttctcattacca
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                                                               TAAGCTGCTTATAACTGGAAATGGCCATTGAGCTGTTTCCTCACAATTGGCGAGATCCCA 1297
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Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M. Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Parsonskis, E., Waterston, R., Williamson, A., Wohldmann, P. and
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Fax: 314 286 1810
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WashU-Merck EST Project
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Location/Qualifiers
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/lab_host="DH10B"
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Pred. No. 0.00e+00;
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Washington University School o
4444 Forest Park Parkway, Box
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1 (bases 1 to 530)

Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,

Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, I

Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F.,

Parevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and
     342;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. High quality sequence stop: 257.

Location/Qualifiers
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                       Similarity
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     Conservative
                                                                                                /tissue_type="Pooled |
pregnant uterus"
/lab_host="DH10B"
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a 96 c 93 g
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/note="Organ: mixed (see below); Vector: pT7T3D-Pac
/note="Organ: mixed (see below); Vector: pt7T3D-Pac
(Pharmacia) with a modified polylinker; Site1: Not I;
Site2: Eco RI; Equal amounts of plasmid DNA from three
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NDHPU, and fetal heart NbHH19W) were mixed, and ss circles
were made in vitro. Following HAP purification, this DNA
                                                                                                                                                                                                                                                reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 260232-265223,
                                                                                                                                                                                                                                                                                                    was used as tracer in a subtractive hybridization
                                                                                                                                                                                                            340488-345479, and 484488-489479."
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EST.
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WashUrHHMI Mouse EST Project
Washington University School of Medicinep
4444 Forest Park Parkway, Box 8501, St. Lo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T., Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
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AA117046
sequence tagged fruit fly.
                                                                     D. melanogaster
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                          g1263754
                                                                                   DM36D7S
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Similarity 76.9%;
30; Conservative
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Vertebrata; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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/lab_host="DH10B"
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/strain="C57BL/6J"
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Gerken, S.C., Matsunami, N., Lawrence, E., Carlson, M., Moore, M., Ballard, L., Melis, R., Robertson, M., Bradley, P., Elsner, T., Tingey, A., Rodriguez, P., Albertsen, H., Lalouel, J.-M. and White, R. Genetic and physical mapping of simple sequence repeat containing sequence tagged sites from the human genome
Unpublished (1993)
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Eukaryotae; mitochondrial eukaryotes; Metazoa; Arthropoda;
Tracheata; Insecta; Pterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                               Submitted by: Utah Center for Human Genome Research University of Utah, Dept. of Human Genetics
2160 Eccles Institute of Human Genetics
Salt Lake City, UT 84112
e-mail: sts@corona.med.utah.edu
primer B: ACCAARICCACGRAGE
Primer B: ACCAARICCACGTCACTTAGA
32P-label: A Primer
PCR Profile:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        g308208
PCR primer; STS sequence; microsatellite marker; repeat; repeat polymorphism; sequence tagged site Homo sapiens DNA.
Homo sapiens
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in_situ_site_primary = 98
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PCR Cycles: 5
Denaturation: 94C 10sec
Annealing: 58C 10sec
Extension: 72C 20sec
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Human STS UT5145.
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database_searched - SWISSPROT
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                                                                              Initial Denaturation: 94C
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/strain="Oregon-R"
/clone="3607"
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23; Mismatches 8;
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Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R. Washu-kerck EST project
Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         zp52d02.rl Stratagene HeLa cell s3 937216 Homo sapiens cDNA clone 613059 5' similar to SW:EPIP_RABIT P10658 PROBABLE PHOSPHOSERINE AAHINOTRANSFERASE ; 61765083
                                                                                                                                                                                                                                                         Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Seq primer: -28Mi3 rev2 from Amersham
High quality sequence stop: 75.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                               Contact: Wilson RK
WashU-Merck EST Project
Washington University Scho
4444 Forest Park Parkway,
                                                                                                                                                                                                                                                                                                                                        Tel: 314 286 1800
Fax: 314 286 1810
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Vertebrata;
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Similarity 70.08;
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/lab_host="SOLR (kanamycin
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                                                                                                                                /clone_lib="Stratagene HeLa cell
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H37487;
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McIntosh L.,
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McIntosh L., Ohlrogge J.,
Retzel E., Somerville C.;
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12-MAR-1997 (Rel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis thaliana (thale cress)
Eukaryotae; mitochondrial eukaryotoes; Viridiplantae;
Charophyta/Embryophyta group; Embryophyta; Magnoliophyta;
Magnoliopsida; Capparales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Plant Physiol. 106:1241-1255(1994).
Contact: Thomas Newman MSU-DOE Plant Research Laboratory Michigan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human STS:
G11923
                                                                                                                                                                                                                                                                                                                                                                                         g1036742
STS sequence; primer; sequence tagged
human STSs derived from random genomic
Contact: Thomas Hudson Whitehead Institute/MIT Center for Genome Research Whitehead Institute for Biomedical Research 9 Cambridge Center, Cambridge MA 02142 USA Tel: 617 252 1900
                                                                                                                                                   Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G11923
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Similarity 74.4%;
32; Conservative
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/Clone="182E10T7"
/strain="var columbia"
/note="thale cress"
P; 72 A; 41 C; 50 G; 70 T;
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51, Last updated, Version 13)
thaliana cDNA clone 182E10T7.
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Raikhel N.,
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Pred. No. 5.
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Email:

thudson@genome.wi.mit.edu

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RESULT 13
LOCUS
DEFINITION
ACCESSION
                               JOURNAL COMMENT
                                                             REFERENCE
AUTHORS
TITLE
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ORIGIN
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Best Local Similarity 70.8%;
Matches 34; Conservative
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G06067
                                                                                          Homo sapiens

Bukaryotae; mitochondrial eukaryotes; Metazoa; Chordata;

Vertebrata; Gnathostomata; Osteichthyes; Sarcopterygil; Cl

Tetrapoda; Amniota; Mammalia; Theria; Eutheria; Archonta;

Catarrhini; Hominidae; Homo.
                                                                                                                                                       g859312
STS sequence; primer; sequence tagged site
human STSs derived from sequences in dbEST
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Contact: Thomas Hudson Whitehead Institute/MIT Center for Whitehead Institute for Biomedical
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                                                             Whitehead Institute/MIT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tris-HCL:
pH: 9.3
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Primer: each 5 pM
dNTPs: each 4 nM
Tag Polymerase: 0.025 units/ul
Total Vol: 20 ul
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Profile:
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KCl: 50 mM
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Annealing: 56 de
Polymerization:
PCR Cycles: 35
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/map="791_B_4; 816_F_6;
963_G_6"
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WI-6406.
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Pred. No. 9.46e-
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Best Local
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Marra M., Hillier L., Allen M., Bowles M., Dietrich N., Dubuque T., Geisel S., Kucaba T., Lacy M., Le M., Martin J Morris M., Schellenberg K., Steptoe M., Tan F., Underwood Moore B., Theising B., Wylie T., Lennon G., Soares B., Wil
                                                                                  Mus musculus (house mouse)
Eukaryotae; mitochondrial eukaryotes; Metazoa;
Vertebrata; Eutheria; Rodentia; Sciurognathi; l
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Primer B: CACAGTTGTGAAGCCGTGC
STS size: 225
PCR Profile:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9 Cambridge Center, Cambridge MA
Tel: 617 252 1900
Fax: 617 252 1902
Email: thudson@genome.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prepared with primer pairs derived Location/Qualifiers
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Similarity 78.4%;
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Primer: each 5 pM
dNTPs: each 4 nM
Tag Polymerase: 0.0
Total Vol: 20 ul
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Annealing: 56 de
Polymerization:
PCR Cycles: 35
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group
group
complement(235..253)
/map-"923_H_8; 959_F_5; 441.7
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29..53
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              Unpublished.

Contact: Marra M/Mouse EST Project WashU-HHMI Mouse EST Project WashIngton University School of MedicineP 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108 Tel: 314 286 1800 Fax: 314 286 1810 Email: mouseest@watson.wustl.edu This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. MGI:401927 Putative full length read vector to vector length is 295 Seq primer: -28m13 rev1 ET from Amersham High quality sequence stop: 272.
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15-MAR-1997 (Rel. 51, Last updated, Version 2)
15-MAR-1997 (Rel. 51, Last updated, Version 2)
mv25008.r1 GuayWoodford Besler mouse kidney day 0 Mus musculus cDNA
clone 656079 5' similar to TR:G497940 G497940 MAJOR VAULT PROTEIN.
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                                                                                                                                                                                                                                                                                                                   Marra M., Hillier L., Allen M., Bowles M., Dietrich N.,
Dubuque T., Geisel S., Kucaba T., Lacy M., Le M., Martin J.,
Morris M., Schellenberg K., Steptoe M., Tan F., Underwood K.,
Moore B., Theising B., Wylie T., Lennon G., Soares B., Wilson
                                                                                                                                                                                                                                                                Waterston R.;
"The WashU-HHMI Mouse EST Project";
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Vertebrata;
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Local Similarity 75.0%;
les 33; Conservation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  musculus (house mouse)
aryotae; mitochondrial eukaryotes; Metazoa;
tebrata; Eutheria; Rodentia; Sciurognathi; j
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/dev_stage="19.5 dpc total fetus"
/dev_stage="19.5 dpc total fetus"
/lab_host="DH10B (ampicillin resistant)"
<1..>267
78 A; 70 C; 61 G; 58 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Minoru
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3003"
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Muridae; M
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completed: Thu ne : 773 secs.
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                                            TEGTCCCCGGAAACCTCAGGGGCTTGGAGGCGGCGG
                                                                                             Similarity
29; Conser
                                                                                                                                              281
                                                                                                                                            BP;
                                                                                              Conservative
                                                                                                                                                                                                                 Guay-Woodford
                                                                                                                                                                                                                       CTCGAGTTTTTTTTTTTTTTT3' Library provided Lisa
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                                                                                                                                           69 A;
                                                                                                                                                                             /clone_lib="GuayWoodford Beier mouse kidney /dev_stage="newborn (day 0)"
                                                                                                                                                                                                                                                                                       /organism="Mus musculus"
/strain="C57BL/6J"
            Aug
                                                                                                                                                                                                      /clone="656079"
                                                                                                                                                                  /lab_host="SOLR (kanamycin resistant)"
                                                                                                        1.4%;
            21 10:48:17 1997
                                                                                                                                           75 C;
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Pred. No. 5.47e-04;
0; Mismatches 7;
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